SEQUENCE LISTING

<110> Salbaum, Michael J.

<120> NOPE Polypeptides, Encoding Nucleic Acids and Methods of Use

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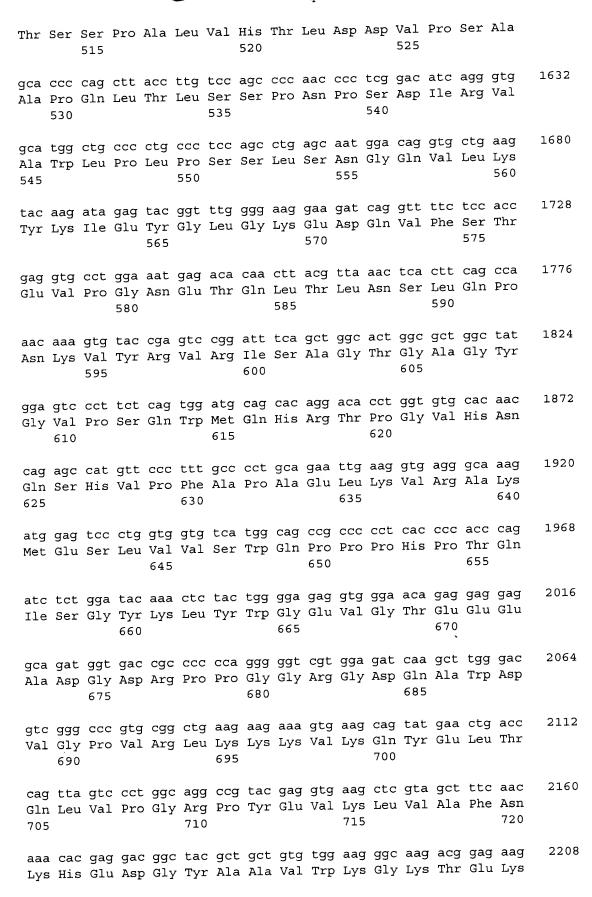
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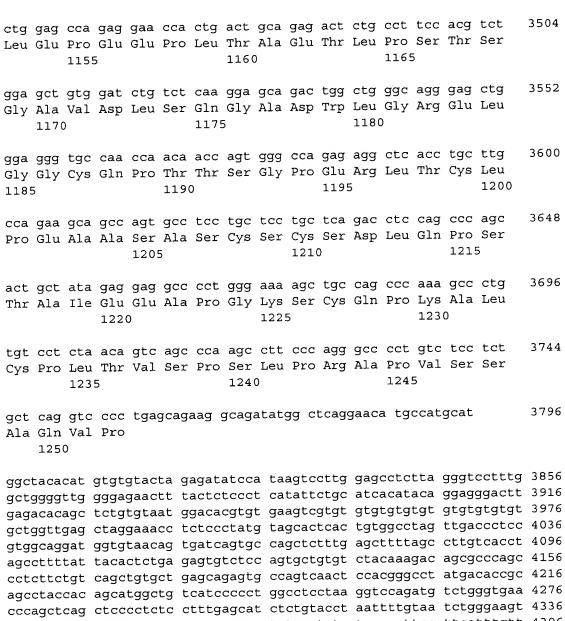
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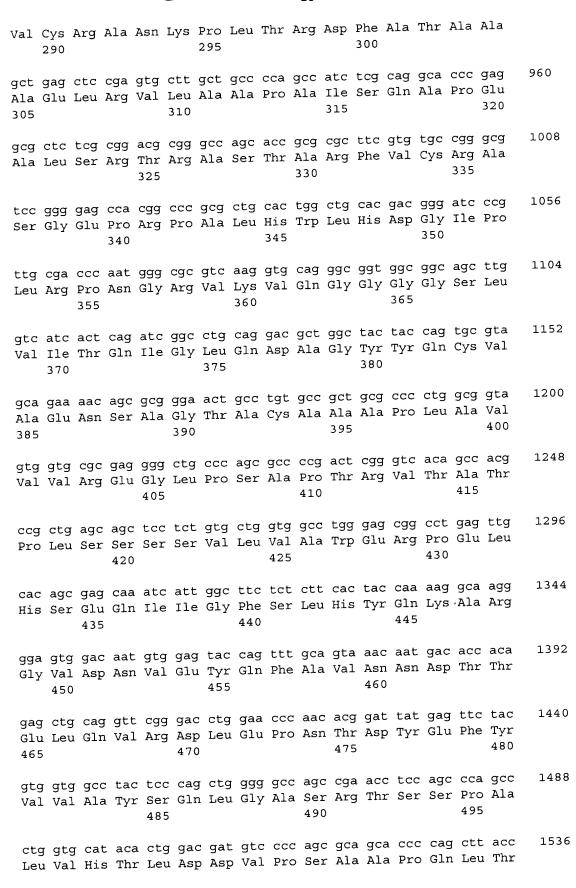
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gaa Glu	cgc Arg	tcc Ser	acc Thr 820	ctg Leu	cct Pro	gac Asp	cgg Arg	cct Pro 825	tca Ser	aca Thr	cct Pro	cct Pro	tct Ser 830	gac Asp	ctg Leu	2496
cgc Arg	ctg Leu	agc Ser 835	ccc Pro	ctg Leu	aca Thr	cca Pro	tcc Ser 840	acc Thr	gtt Val	cgg Arg	tta Leu	cac His 845	tgg Trp	tgt Cys	ccc Pro	2544
ccc Pro	acg Thr 850	gag Glu	ccc Pro	aat Asn	ggt Gly	gag Glu 855	att Ile	gtg Val	gag Glu	tat Tyr	cta Leu 860	att Ile	ctc Leu	tac Tyr	agc Ser	2592
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gga Gly	aac Asn	atc Ile	ttc Phe	agt Ser 885	Ala	gag Glu	gtc Val	cat His	ggc Gly 890	cta Leu	gag Glu	agt Ser	gac Asp	act Thr 895	cgg Arg	2688
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325

Ser Gly Glu Pro Arg Pro Ala Leu His Trp Leu His Asp Gly Ile Pro 345 Leu Arg Pro Asn Gly Arg Val Lys Val Gln Gly Gly Gly Ser Leu 360 355 Val Ile Thr Gln Ile Gly Leu Gln Asp Ala Gly Tyr Tyr Gln Cys Val 375 Ala Glu Asn Ser Ala Gly Thr Ala Cys Ala Ala Ala Pro Leu Ala Val 395 390 Val Val Arg Glu Gly Leu Pro Ser Ala Pro Thr Arg Val Thr Ala Thr 410 405 Pro Leu Ser Ser Ser Val Leu Val Ala Trp Glu Arg Pro Glu Leu 425 His Ser Glu Gln Ile Ile Gly Phe Ser Leu His Tyr Gln Lys Ala Arg 440 Gly Val Asp Asn Val Glu Tyr Gln Phe Ala Val Asn Asn Asp Thr Thr 460 455 Glu Leu Gln Val Arg Asp Leu Glu Pro Asn Thr Asp Tyr Glu Phe Tyr 475 470 Val Val Ala Tyr Ser Gln Leu Gly Ala Ser Arg Thr Ser Ser Pro Ala 485 Leu Val His Thr Leu Asp Asp Val Pro Ser Ala Ala Pro Gln Leu Thr 505 Leu Ser Ser Pro Asn Pro Ser Asp Ile Arg Val Ala Trp Leu Pro Leu 520 515 Pro Ser Ser Leu Ser Asn Gly Gln Val Leu Lys Tyr Lys Ile Glu Tyr 540 535 Gly Leu Gly Lys Glu Asp Gln Val Phe Ser Thr Glu Val Pro Gly Asn 555 550 Glu Thr Gln Leu Thr Leu Asn Ser Leu Gln Pro Asn Lys Val Tyr Arg 570 565 Val Arg Ile Ser Ala Gly Thr Gly Ala Gly Tyr Gly Val Pro Ser Gln 585 580 Trp Met Gln His Arg Thr Pro Gly Val His Asn Gln Ser His Val Pro 600 Phe Ala Pro Ala Glu Leu Lys Val Arg Ala Lys Met Glu Ser Leu Val 615 Val Ser Trp Gln Pro Pro Pro His Pro Thr Gln Ile Ser Gly Tyr Lys 635 630 Leu Tyr Trp Gly Glu Val Gly Thr Glu Glu Glu Ala Asp Gly Asp Arg 650 645 Pro Pro Gly Gly Arg Gly Asp Gln Ala Trp Asp Val Gly Pro Val Arg 665 Leu Lys Lys Lys Val Lys Gln Tyr Glu Leu Thr Gln Leu Val Pro Gly 685 680 Arg Pro Tyr Glu Val Lys Leu Val Ala Phe Asn Lys His Glu Asp Gly 695 Tyr Ala Ala Val Trp Lys Gly Lys Thr Glu Lys Ala Pro Thr Pro Asp 715 710 Leu Pro Ile Gln Arg Gly Pro Pro Leu Pro Pro Ala His Val His Ala 730 725 Glu Ser Asn Ser Ser Thr Ser Ile Trp Leu Arg Trp Lys Lys Pro Asp 745 Phe Thr Thr Val Lys Ile Val Asn Tyr Thr Val Arg Phe Gly Pro Trp

755 760 Gly Leu Arg Asn Ala Ser Leu Val Thr Tyr Tyr Thr Ser Ser Gly Glu 780 775 Asp Ile Leu Ile Gly Gly Leu Lys Pro Phe Thr Lys Tyr Glu Phe Ala 795 790 Val Gln Ser His Gly Val Asp Met Asp Gly Pro Phe Gly Ser Val Val 810 805 Glu Arg Ser Thr Leu Pro Asp Arg Pro Ser Thr Pro Pro Ser Asp Leu 825 Arg Leu Ser Pro Leu Thr Pro Ser Thr Val Arg Leu His Trp Cys Pro 845 840 Pro Thr Glu Pro Asn Gly Glu Ile Val Glu Tyr Leu Ile Leu Tyr Ser 855 Asn Asn His Thr Gln Pro Glu His Gln Trp Thr Leu Leu Thr Thr Glu 875 Gly Asn Ile Phe Ser Ala Glu Val His Gly Leu Glu Ser Asp Thr Arg 890 885 Tyr Phe Phe Lys Met Gly Ala Arg Thr Glu Val Gly Pro Gly Pro Phe 905 900 Ser Arg Leu Gln Asp Val Ile Thr Leu Gln Glu Thr Phe Ser Asp Ser 925 920 915 Leu Asp Val His 930 <210> 5 <211> 825 <212> DNA <213> Mus musculus <220> <221> CDS <222> (1)...(825) <400> 5 cga caa agc tcc cac agg gaa gcc ctt ccc gga ttg tcc tcc tca ggc 48 Arg Gln Ser Ser His Arg Glu Ala Leu Pro Gly Leu Ser Ser Ser Gly acc cca gga aac cca gcg ctc tac aca aga gct cgg ctt ggg cct ccc Thr Pro Gly Asn Pro Ala Leu Tyr Thr Arg Ala Arg Leu Gly Pro Pro 25 20 agt gtc cct gct gcc cat gag ttg gag tcc ctc gtg cat cct cgt ccc 144 Ser Val Pro Ala Ala His Glu Leu Glu Ser Leu Val His Pro Arg Pro 35 cag gat tgg tcc cca cca ccc tca gat gtg gaa gac aag gct gaa gta 192 Gln Asp Trp Ser Pro Pro Pro Ser Asp Val Glu Asp Lys Ala Glu Val 55 cac age ctt atg ggt ggc agt gtt tca gat tgc cgg ggc cac tcc aag His Ser Leu Met Gly Gly Ser Val Ser Asp Cys Arg Gly His Ser Lys

105

120

7.0

65

245

260

cag gtc ccc Gln Val Pro

275

Pro Leu Thr Val Ser Pro Ser Leu Pro Arg Ala Pro Val Ser Ser Ala

265

100

115

80 75 aga aag atc tcc tgg gct cag gca ggg gga cca aac tgg gca ggc tcc Arg Lys Ile Ser Trp Ala Gln Ala Gly Gly Pro Asn Trp Ala Gly Ser tgg gca ggc tgt gag ctg ccc cag ggt agt ggt cca agg ccg gct ctg 336 Trp Ala Gly Cys Glu Leu Pro Gln Gly Ser Gly Pro Arg Pro Ala Leu acc cgt gct ctg ctg cct cca gcg gga acc ggg cag aca ctg ctg ctg 384 Thr Arg Ala Leu Leu Pro Pro Ala Gly Thr Gly Gln Thr Leu Leu Leu 432 480 155 528 175 576 190 624 672 720 240 235 Ala Ile Glu Glu Ala Pro Gly Lys Ser Cys Gln Pro Lys Ala Leu Cys 255 cet cta aca gtc agc cca agc ctt ccc agg gcc cct gtc tcc tct gct 816

270

825

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Gly aaa	cct Pro	ccg Pro	acc Thr 20	agg Arg	gtg Val	aca Thr	tgg Trp	agc Ser 25	aag Lys	gat Asp	gga Gly	gac Asp	act Thr 30	gta Val	cta Leu	96
gag Glu	cat His	gag Glu 35	aac Asn	ctg Leu	cac His	ctg Leu	cta Leu 40	ccc Pro	aat Asn	ggc	tcc Ser	ctg Leu 45	tgg Trp	ctg Leu	tcc Ser	144
tca Ser	ccc Pro 50	cta Leu	gag Glu	caa Gln	gaa Glu	gac Asp 55	agc Ser	gat Asp	gat Asp	gag Glu	gaa Glu 60	gct Ala	ctt Leu	agg Arg	atc Ile	192
tgg Trp 65	aag Lys	gtc Val	act Thr	gag Glu	ggc Gly 70	agc Ser	tat Tyr	tcc Ser	tgt Cys	ctg Leu 75	gcc Ala	cac His	agc Ser	ccg Pro	cta Leu 80	240
gga Gly																243

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ggc ctg cag gac gct ggc tac tac cag tgc gta gca gaa aac agc gcg

192

Gly Leu Gln Asp Ala Gly Tyr Tyr Gln Cys Val Ala Glu Asn Ser Ala

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55

50

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 Pro
 Ala
 Glu
 Leu
 Lys
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 Arg
 Ala
 Lys
 Met
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 Ser
 Leu
 Val

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 10
 10
 15
 15

 Val
 Ser
 Trp
 Gln
 Pro
 Pro
 Pro
 Thr
 Gln
 Thr
 Gln
 Ile
 Ser
 Gly
 Tyr
 Lys

 Leu
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 Trp
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 Asp
 Gly
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 Glu
 Asp
 Gly
 Arg

 Pro
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 Gly
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 Gly
 Asp
 Gln
 Ala
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 Pro
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Trp Leu Arg Trp Lys Lys Pro Asp Phe Thr Thr Val Lys Ile Val Asn
20 25 30

tac act gta cgc ttc ggc ccc tgg ggg ctc agg aat gct tcc ctg gtc 144 Tyr Thr Val Arg Phe Gly Pro Trp Gly Leu Arg Asn Ala Ser Leu Val

Maria Maria

Beng abap ham

1

. . .

12

[]

25

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aca Thr	ctg Leu 50	ctc Leu	acc Thr	aca Thr	gag Glu	gga Gly 55	aac Asn	atc Ile	ttc Phe	agt Ser	gca Ala 60	gag Glu	gtc Val	cat His	ggc Gly	192
cta Leu 65	gag Glu	agt Ser	gac Asp	act Thr	cgg Arg 70	tat Tyr	ttc Phe	ttc Phe	aag Lys	atg Met 75	gga Gly	gcc Ala	cgc Arg	aca Thr	gag Glu 80	240
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